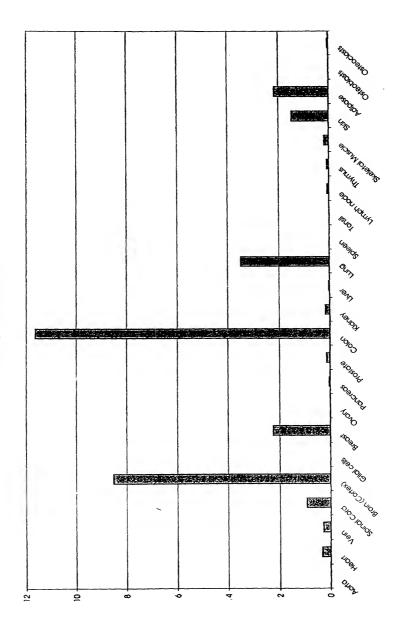
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GAA ATT TCG ATT TCC ACA CTG ATT TCG CCC ACC ATT CCT GGA GAG ATC TCG TGG GAT GTC 627 TLNFLVPGLVI TCT TTT GTT ACT TTG AAC TTC TTG GTG CCA GGA CTG GTC ATT GTG ATC AGT TAC TCC AAA 687 ILQITKASRKRLT ATT TTA CAG ATC ACA AAG GCA TCA AGG AAG AGG CTC ACG GTA AGC CTG GCC TAC TCG GAG 747 V S Q Q D F R L F R AGO CAG CAG ATC CGC GTG TCC CAG CAG GAC TTC CGG CTC TTC CGC ACC CTC TTC CTC CTC 807 F F I M W S P I I I T I L L I ATG GTC TCC TTC TTC ATC ATG TGG AGC CCC ATC ATC ATC ATC ATC CTC CTC ATC CTG ATC 867 Q N F K Q D L V I W P S .L F F W V V A F 309 CAG AAC TTC AAG CAA GAC CTG GTC ATC TGG CCG TCC CTC TTC TGC GTG GTG GCC TTC 927 Figure 1B

## PECAQTT, G CGCCATCTTCCCGGACGCGTGGGCCGGGCGCCCGGC ATG TCC CCT GAG TGT GCA CAG ACG ACG GGC PSHTLDQVVNRTHFPFFFSD CCT GGT CCC TCG CAC ACC CTG GAC CAA GTC AAT CGC ACC CAC TTC CCT TTC TCG GAT 90 S E T 50 GTC AAG GGC GAC CAC CGG TTG GTG TTG AGC GTC GTG GAG ACC ACC GTT CTG GGA CTC ATC 150 N TIT GTC GTC TCA CTG CTG GGC AAC GTG TGT GCT CTA GTG CTG GTG GGC CGC CGT CGG CGC 210 90 I. L 'N T. C A D CGT GGG GCG TCA GCC AGC CTG GTG CTC AAC CTC TTC TGC GCG GAT TTG CTC TTC ACC AGC 270 R W T 110 GCC ATC CCT CTA GTG CTC GTG CGC TGG ACT GAG GCC TGG CTG TTG GGG CCC GTC GTC M T M S G S V T TGC CAC CTG CTC TTC TAC GTG ATG ACA ATG AGC GGC AGC GTC ACG ATC CTC ACA CTG GCC 390 ν CIVRL 150 R M R R GCG GTC AGC CTG GAG CGC ATG GTG TGC ATC GTG CGC CTC CGG CGC GGC TTG AGC GGC CCG 450 170 GGG CGG CGG ACT CAG GCG GCA CTG CTG GCT TTC ATA TGG GGT TAC TCG GCG CTC GCC GCG 510 190 CTG CCC CTC TAC ATC TTG TTC CGC GTG GTC CCG CAG CGC CTT CCC GGC GGG GAC CAG GAA 570 210 T L N R I G E I S W ATT CCG ATT TGC ACA TTG GAT TGG CCC AAC CGC ATA GGA GAA ATC TCA TGG GAT GTG TTT 630 F v P G T. т т 230 TTT GAG ACT TTG AAC TTC CTG GTG CCG GGA CTG GTC ATT GTG ATC AGT TAC TCC AAA ATT 690 250 TTA CAG ATC ACG AAA GCA TCG CGG AAG AGG CTT ACG CTG AGC TTG GCA TAC TCT GAG AGC 750 DYRLFRTL 270 CAC CAG ATC CGA GTG TCC CAA CAA GAC TAC CGA CTC TTC CGC ACG CTC TTC CTG CTC ATG 810 290 ITI S P · T τ GTT TCC TTC ATC ATG TGG AGT CCC ATC ATC ATC ATC CTC CTC ATC TTG ATC CAA 870 310 AAC TTC CGG CAG GAC CTG GTC ATC TGG CCA TCC CTT TTC TTC TGG GTG GTG GCC TTC ACG 930 330 s TTT GCC AAC TCT GCC CTA AAC CCC ATA CTG TAC AAC ATG TCG CTG TTC AGG AAC GAA TGG 990 P E K G 350 AGG AAG ATT TTT TGC TGC TTC TTT TTT CCA GAG AAG GGA GCC ATT TTT ACA GAT AGG TCT 1050 362 GTC AGG CGA AAT GAC TTG TCT GTT ATT TCC AGC TAA 108€ CTAGCCTCTGGTGCCAGGTGAACCACGGTGTGCATGTAAAGGGAGTTAACTTCAAGGAAAGCCCACCAGTGCCCCCTGC TTTAAAAATACCCGACTTCCAACAGCAGGCATCTACGGAGCCAGCAAATTAAGGAATGATCGCTCAGTATAAAAATATT

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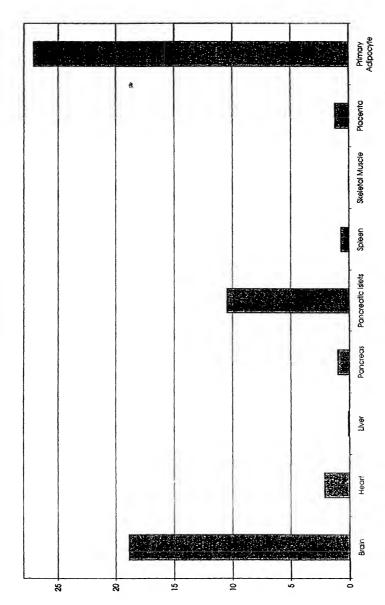
10035151 CTCCCT

Expression of 14273 in Human Tissues Pathology Phase I (subset)

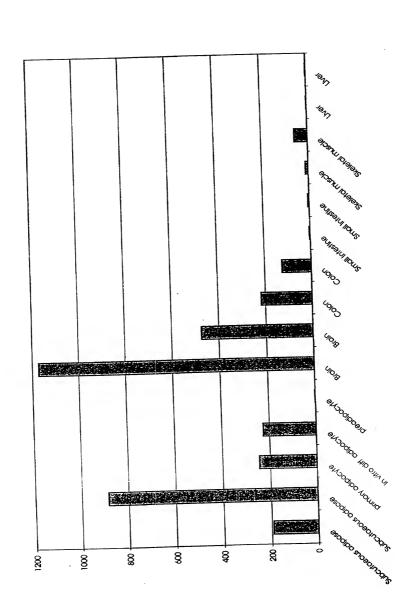


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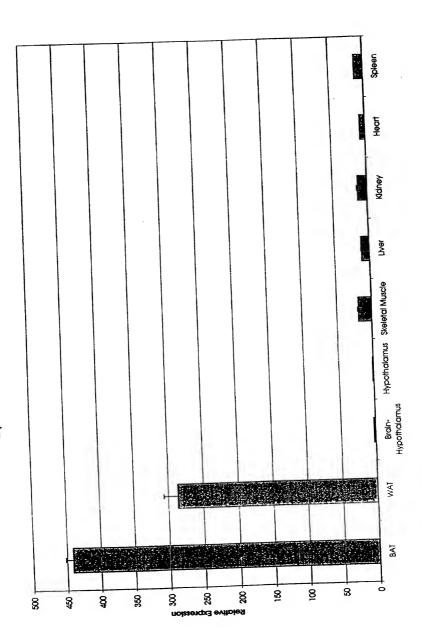
Expression of 14273 in Human Tissues Metabolic Phase I



Expression of 14273 in Human Tissues Metabolic Phase II



Expression of 14273 in Normal Mouse Tissues



## **Expression of 14273 in Normal Mouse Tissues**

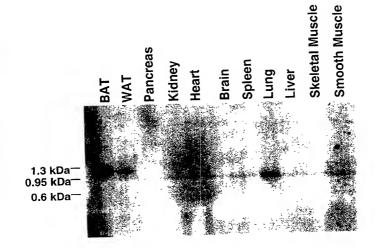
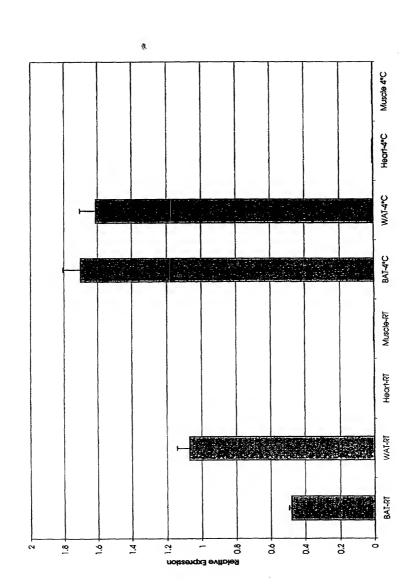


Figure 5

Regulation of 14273 Expression by Cold Exposure



The first and th

WAT-agoutt Regulation of 14273 Expression in Genetic Models of Obesity WAT-ob/ob WAT-wildtype 20 8 8 3 8 8 8 Relative Expression

## BW on High Fat Diet (male)

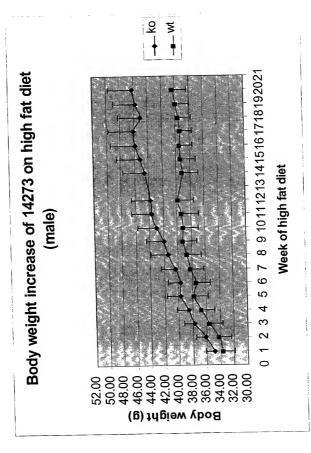


FIGURE 7